EXHIBIT B

```
<110>   Hanski, Emanuel
        Moses, Allon E
        Hidalgo-Grass, Carlos

<120>   Compositions and methods for the treatment and prophylaxis of
        infections caused by gram positive bacteria

<130>   73975/JPW/JW

<140>   US 10/525,178
<141>   2005-02-22

<150>   IL 151436
<151>   2002-08-22

<150>   PCT/IL03/00687
<151>   2003-08-19

<160>   32

<170>   PatentIn version 3.3

<210>   1
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<220>
<223>   m13/puc sequence primer (-20)

<400>   1
gtaaaaaacg acggccagt                                                  19


<210>   2
<211>   16
<212>   DNA
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<220>
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        amplification

<400>   2
aacagctatg accatg                                                     16


<210>   3
<211>   20
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<220>
<223>   Reverse primer for tag amplification

<400>   3
agcagttcgt agttatcttg                                                 20


<210>   4
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<212> DNA
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<400> 4
ttatcagcaa taaaccagc   19

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aaagtcctcc tgggtatg   18

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<400> 6
tttggcagct ttgacgatgc   20

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<220>
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<400> 7
tcttcaagca gctgattggg   20

<210> 8
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<220>
<223> 2598-2620 in sil

<400> 8
ggagttggtt tatcaaatgt cag   23

<210> 9
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<223>   3213-3235 in sil

<400>   9
atctgccaca aagactgatc aag                                    23


<210>   10
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<220>
<223>   2013-2033 in sil

<400>   10
ttattggatc ggaacttacg c                                      21


<210>   11
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<220>
<223>   3554-3574 in sil

<400>   11
tgcttcccaa caacttacca c                                      21


<210>   12
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<220>
<223>   2088-2109 in sil

<400>   12
gctcgctata gtaagcaaat cg                                     22


<210>   13
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<220>
<223>   5871-5888 in sil

<400>   13
cagcgattaa gcattgac                                          18


<210>   14
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<220>
<223>   1616-1634 in sil
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<400>  14
acgaaaggtc aatggttcac                                            20


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<220>
<223>  2338-2357 in sil

<400>  15
aggtatggat aagcgttgag                                            20


<210>  16
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<220>
<223>  3873-3894 in sil

<400>  16
atgacacttg ttacacgtcc                                            20


<210>  17
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<223>  3873-3984

<400>  17
actagtcagc ttgacgaact tc                                         22


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<400>  18
tattcgctta gaaaattaa                                             19


<210>  19
<211>  20
<212>  DNA
<213>  Artificial Sequence

<220>
<223>  emm typing reverse primer

<400>  19
gcaagttctt cagcttgttt                                            20
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<223>  aad9 forward primer

<400>  20
ccatggtcct cgagctctag atcttaag                                    28


<210>  21
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<212>  DNA
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<223>  aad9 reverse primer

<400>  21
ctgcaggcgc ttaccaatta gaatg                                       25


<210>  22
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<212>  DNA
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<220>
<223>  6873-6896 in JS95 sil, 5096-5119 in M1

<400>  22
tcgatatgga gataaagaaa ctgg                                        24


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<220>
<223>  6804-6825 in M1 section 36

<400>  23
aacagtgctt tcaggaactc ct                                          22


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<223>  10031-10052 in M1 section 36

<400>  24
ctaggtgcaa ttgaggagtc aa                                          22


<210>  25
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<220>
<223>  20-43 in JS95 sil, 7287-7306 section 152 in M1

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tcctcgcact gttccaatag                                              20


<210>  26
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<220>
<223>  3580-3599 in M1 section 36

<400>  26
aggtggtgtt ggagcaggta                                              20


<210>  27
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<220>
<223>  1545-1565 in M1 section 36

<400>  27
aagaagtggt cccaatttct g                                            21


<210>  28
<211>  30
<212>  DNA
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<220>
<223>  Forward all M primer with BamHI site

<400>  28
cctgaaaatg aggatccttc ctaaaaaacg                                   30


<210>  29
<211>  32
<212>  DNA
<213>  Artificial Sequence

<220>
<223>  Reverse all M p[rimer with PstI site

<400>  29
gggggctgca gagcttagtt ttcttctttg cg                                32


<210>  30
<211>  20
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<220>
<223>  3' of mga

<400>  30
gattccagaa gcgattattg                                                    20


<210>  31
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<220>
<223>  Leader peptide region of scpA

<400>  31
aatggcaagt ttatcaaatg g                                                  21


<210>  32
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<212>  PRT
<213>  Artificial Sequence

<220>
<223>  SilCR peptide

<400>  32

Asp Ile Phe Lys Leu Val Ile Asp His Ile Ser Met Lys Ala Arg Lys
1               5                   10                  15

Lys
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